#### Femtoliter Injection of ESCRT-III Proteins into Giant Unilamellar Vesicles

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16 [Abstract] The endosomal sorting complex required for transport (ESCRT) machinery mediates 17 membrane fission reactions that exhibit a different topology from that observed in clathrin-coated vesicles. In all of the ESCRT-mediated events, the nascent vesicle buds away from the cytosol. However, 18 19 ESCRT proteins are able to act in membranes with different geometries. For instance, the formation of 20 multivesicular bodies (MVBs) and the biogenesis of extracellular vesicles, both of which require the 21 participation of the ESCRT-III sub-complex, differ in the initial membrane geometry before budding starts 22 and whether the protein complex acts from outside the membrane organelle (causing inward budding) 23 of from within (causing outward budding). Several studies have reconstituted the action of the ESCRT-24 III subunits in supported bilayers and cell-sized vesicles mimicking the geometry occurring during MVBs 25 formation (in-bud), however extracellular vesicle budding (out-bud) mechanisms remain less explored 26 because of the outstanding difficulties of encapsulation of functional ESCRT-III in vesicles. Here, we 27 provide a different approach that allows the recreation of the out-bud formation by combining giant 28 unilamellar vesicles as a membrane model and a microinjection system. The vesicles are immobilized 29 prior to injection via weak adhesion to the chamber coverslip, which also ensures preserving the membrane excess area required for budding. After protein injection, vesicles exhibit outward budding. 30 31 The approach presented in this work can be used in the future to disentangle the mechanisms underlying ESCRT-III-mediated fission recreating the geometry of extracellular bud production, which remains a 32 challenge in spite of all the efforts made to understand it. Moreover, the microinjection methodology can 33 be also adapted to interrogating the action of other cytosolic components on the encapsulating 34 35 membranous organelle. 36

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#### 40 **Graphic abstract**:



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Keywords: Giant unilamellar vesicle (GUV), Microinjection, ESCRT-III, Extracellular vesicles, Adhesion,
budding

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[Background] Extracellular vesicles (EVs) are defined as cell-derived vesicles confined by a lipid bilayer. 46 47 They participate in cellular disposal and in intercellular communication by delivering antigens, genetic material and lipids to recipient cells (Raposo and Stahl, 2019). EV secretion seems to be an ubiquitous 48 process present throughout all kingdoms of life and, in most of the cases, it is related to normal 49 50 physiological conditions (Herrmann et al., 2021). However, a considerable increase in EV number is 51 observed during pathogenic processes including cancer (Clos-Garcia et al., 2018; Tao et al., 2019; 52 Wolfers et al., 2001; Xu et al., 2018), hereditary α-tryptasemia (Glover et al., 2019), multiple sclerosis 53 (Moyano et al., 2016), cytotoxic-drug intoxication (Keklikoglou et al., 2019), and parasitic diseases, 54 among which cerebral malaria has been widely characterized (Combes et al., 2004; Schofield and Grau, 2005; Campos et al., 2010; Mfonkeu et al., 2010; Nantakomol et al., 2011; Martin-Jaular et al., 2011). 55 56 EVs can be classified into two major classes depending on their size and origin: exosomes with a typical

57 diameter of 30-150 nm, and microvesicles that have a diameter of 100-1000 nm (Raposo and Stahl, 58 2019). Whereas microvesicles are shed by outward budding of the plasma membrane, exosomes are 59 generated by the fusion of multivesicular bodies (MVBs) with the plasma membrane followed by the release of intraluminal vesicles (ILVs) (van Niel et al., 2018). EV populations are considered highly 60 61 heterogeneous both in content and in size, probably due to the different pathways along which they 62 originate from (Raposo and Stahl, 2019), making it difficult to resolve the associated mechanisms of 63 cargo-packing and specific activity during cell signaling. Understanding the molecular processes that govern their biogenesis via employing simple mimetic systems could provide a clue to solve the 64 65 mechanism of action, and therefore, help to understand the pathophysiology of certain diseases.

66 As mentioned above, EVs can be generated by different pathways. Classically, exosomes are

generated from the endosomal system as ILVs in the MVBs, which represent late endosomes that have 67 68 suffered inward budding of their membrane. This process is orchestrated by the endosomal sorting 69 complex required for transport (ESCRT) machinery, which comprises several protein subunits organized into four different complexes (ESCRT-0, -I, -II and -III) and the accessory Vps4 complex (reviewed in 70 71 (Vietri et al., 2020)). Typically, ESCRT complexes are recruited to the endosomal membrane in a 72 stepwise manner. The process begins with recognition of monoubiguitinated cargo by ESCRT-0 73 (Raiborg and Stenmark, 2009). Then, ESCRT-0 recruits ESCRT-I to the endosomal membrane 74 (Katzmann et al., 2003). ESCRT-I activates ESCRT-II producing the membrane invagination (Gill et al., 75 2007), which in turn activates ESCRT-III (Vps2, Vps20, Vps24 and Vps32) assembly, a process required 76 for ILV scission into the MVB lumen (Babst et al., 2002; Im et al., 2009; Teis et al., 2010). Subsequently, 77 Vps4 AAA ATPase catalyzes the dissociation and recycling of ESCRT-III components from the 78 membrane (Obita et al., 2007; Lata et al., 2008), which cooperatively drive membrane scission 79 (Chiaruttini et al., 2015; Mierzwa et al., 2017; Schöneberg et al., 2018). In other cases, there is a parallel 80 way to recruit ESCRTs to endosomal membranes. Two non-canonical pathways have been identified so far: 1) activation of ESCRT-I by Bro1, which functions as ubiquitin acceptor (Tang et al., 2016), and 2) 81 82 ESCRT-III activation by Alix, which mediates the ubiquitin-independent, but ESCRT-III-dependent endocytosis (Dores et al., 2012). On the other hand, the ESCRT machinery has also been implicated in 83 84 the production of nano-sized vesicles that are enriched in cell surface proteins, reflecting its participation 85 during microvesicle formation (Nabhan et al., 2012; Wang and Lu, 2017).

Plasmodium falciparum and other strictly intracellular protozoans are devoid of ESCRT-0. -I and -II 86 87 complexes. However, our previous results demonstrated that there is a minimal and functional ESCRT-III machinery present in *P. falciparum* in which PfBro1 activates PfVps32 and PfVps60, both ESCRT-III 88 89 members, triggering EV biogenesis (Avalos-Padilla et al., 2021b). Involving an intracellular parasite, the 90 study of this process is problematic. Moreover, the knockdown or deletion of ESCRT genes in other 91 organisms results in the formation of aberrant structures that lack ILVs (Doyotte et al., 2005; Nickerson 92 et al., 2006). To address these difficulties, membrane models have been widely used to analyze in vitro 93 ESCRT-III-mediated events. In this direction, giant unilamellar vesicles (GUVs) (R. Dimova and Marques, 94 2019; Rumiana Dimova, 2019) combined with ESCRT proteins have become an established platform to 95 examine the formation of MVBs (Im et al., 2009; Avalos-Padilla et al., 2018; Booth et al., 2019; Avalos-96 Padilla et al., 2021a; Algabandi et al., 2021). To mimic the geometry occurring in this process, ESCRT 97 components are introduced in the vesicle surroundings. The proteins induce membrane invaginations 98 towards the vesicle interior, which can lead to the formation of ILVs connected to the mother membrane 99 through a thin neck and the final cleavage of the neck results in the formation of MVB-like GUVs.

The out-budding processes (as in the formation of microvesicles shed by the plasma membrane) exhibit a reverse budding topology, compared to that of MVB formation. Thus, to explore such process the ESCRT units have to act from within the vesicle model. In other words, the proteins have to be introduced into the GUVs lumen. One approach to accomplish this consists of ESCRT-proteins encapsulation inside GUVs by forming the vesicles in the presence of the proteins (Schöneberg *et al.*, 2018). However, under this condition, one cannot observe the vesicle response during and immediately

# bio-protocol

after introducing the proteins. To evade these drawbacks, we have designed an approach in which preformed GUVs encapsulating the buffer necessary for protein activity are injected with the ESCRT-III proteins. With this technique we are able to observe in real time the dynamics of out-bud formation (mimicking the process driven during EV biogenesis), and to evaluate the effects specific to a particular protein.

111 Injection approaches in GUVs have been applied previously (Wick et al., 1996; Hurtig and Orwar, 112 2008; Lefrançois et al., 2018). In isolated GUVs, it is important to ensure control over the vesicle volume and area. In particular, the injection if isotonic solutions can pull out the excess membrane area needed 113 114 for deformation, which would then prohibit outward budding. We thus adapted the protocol employing osmolarities of the injected solutions which lead to vesicle deflation to allow for creating excess vesicle 115 116 area for deformation and budding. Furthermore, isolated vesicles need to be immobilized to facilitate 117 the puncturing of the membrane without displacing and losing them. In previous work, the immobilization was ensured by working with GUVs which are directly formed on the substrate for GUV swelling. 118 119 However, such vesicles are typically connected to other GUVs and structures, thus not ensuring 120 area/volume conservation. Here, we employed biotin-avidin-based adhesion, using biotinylated lipids in 121 the vesicle and an avidin-coated substrate to which the GUVs were fixed as proposed earlier (Maan et al., 2018). Successful injection of ESCRTs and further registration of the functionality of the proteins, 122 123 namely formation of out-buds, requires fine adjustment of the adhesion level. On one hand, strong 124 adhesion stabilizes the vesicles during the puncturing procedure, while on the other hand, it increases the membrane tension while consuming the area available for deformation. The latter effect limits the 125 126 ability of the membrane to bend and thus hinders budding. We thus optimized the avidin surface concentration to ensure mild adhesion. Another important aspect to consider is the buffer in which 127 128 proteins remain active, in the case of ESCRT-III proteins, the used buffer is 150 mM NaCl and 25 mM 129 Tris-HCl, pH 7.4 (~ 325 mOsmol/Kg). This high salt concentration hampers the growing of GUVs by the 130 standard electroformation protocol (Angelova and Dimitrov, 1986); therefore, we used the gel-assisted 131 method (Weinberger et al., 2013) in which we were able to grow vesicles encapsulating this buffer. As we are working with P. falciparum ESCRT-III subunits, the GUV lipid composition was selected to mimic 132 the inner leaflet of the red blood cell plasma membrane (Virtanen et al., 1998), however, this can be 133 134 modified depending on the system. We demonstrate the injection and outward budding process also for 135 GUVs injected with another ESCRT-III system, namely the protozoan parasite responsible for amoebiasis, Entamoeba histolytica, whose characterization in GUVs has been previously reported 136 137 (Avalos-Padilla et al., 2018), and using a suitable membrane composition. Finally, as mentioned above, to deflate the GUVs and thus to ensure that the vesicles exhibit excess membrane area, needed for the 138 139 formation of the out-buds, the injected proteins were kept in a 0.8× buffer. As a control, upon injection of GUVs with the same volume and osmolarity of protein-free solution as in the experimental set-up, no 140 141 out-buds appeared, demonstrating the validity of our approach. 142

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145	<u>Ma</u>	Materials and Reagents			
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147	Α.	Lipi	ids		
148		1.	1-palmitoyl-2-oleoyl- <i>sn</i> -glycero-3-phosphocholine [POPC] (Avanti Polar Lipids, catalog number:		
149			850457)		
150 151		2.	1-palmitoyl-2-oleoyl- <i>sn</i> -glycero-3-phospho-L-serine [POPS] (Avanti Polar Lipids, catalog number: 840034)		
152		3.	Cholesterol (ovine) [chol] (Avanti Polar Lipids, catalog number: 700000)		
153		4.	1,2-dioleoyl- <i>sn</i> -glycero-3-phospho-(1'-myo-inositol-3'-phosphate) [PI(3)P] (Avanti Polar Lipids,		
154			catalog number: 850150)		
155		5.	1,2-distearoyl- <i>sn</i> -glycero-3-phosphoethanolamine-N-[biotinyl(polyethyleneglycol)-2000]		
156			[DSPE-PEG 2000 Biotin] (Avanti Polar Lipids, catalog number: 880129)		
157		6.	1,2-dipalmitoyl-sn-glycero-3-phosphoethanolamine-N-(lissamine rhodamine B sulfonyl) [DPPE-		
158			Rhodamine] (Avanti Polar Lipids, catalog number: 810158)		
159		Not	te: All lipids were dissolved in chloroform at a final concentration of 10 mg/mL and stored at -		
160		20	°C for up to one month.		
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162	В.	Rea	agents		
163		1.	Avidin from egg white (Sigma-Aldrich, catalog number: A9275)		
164		2.	Bovine Serum Albumin [BSA] (Sigma-Aldrich, catalog number: A8806)		
165		3.	Biotinylated BSA (Thermo Fisher Scientific, catalog number: 29130)		
166		4.	Chloroform (Merck, Supelco, catalog number: 288306)		
167		5.	Ethanol absolute (Merk, catalog number: 1009831011)		
168		6.	Methanol (Merk, catalog number: 34860)		
169		7.	Milli-Q <sup>®</sup> water (Millipore system)		
170		8.	Polyvinyl alcohol [PVA] (Approximate MW 145, 000 g/mol, Merck, catalog number: 814894)		
171		9.	Polyethylene glycol fluorescein isothiocyanate [PEG-FITC] (Nanocs, catalog number: PG1-FC-		
172			40k)		
173		10.	NaCl (Sigma-Aldrich, catalog number: S7653)		
174		11.	Trizma <sup>®</sup> hydrochloride (Sigma-Aldrich, catalog number: T6666)		
175		12.	Purified recombinant PfBro1 and PfVps32 (Avalos-Padilla et al., 2021b)		
176		13.	Purified recombinant EhVps20t and EhVps32 (Avalos-Padilla et al., 2018)		
177		14.	PVA solution (5% w/v) (see Recipe 1)		
178		15.	Lipid mix (1 mg/mL) for the injection of ESCRTs purified from <i>P. falciparum</i> containing		
179			POPC:POPS:DSPE-PEG 2000 Biotin:DPPE-Rhodamine (see Recipe 2)		
180		16.	Lipid mix (1 mg/mL) for the injection of ESCRTs purified from <i>Entamoeba histolytica</i> containing		
181			POPC:POPS:chol:PI(3)P:DSPE-PEG 2000 Biotin:DPPE-Rhodamine (see Recipe 3)		
182		17.	Protein buffer 1×, pH 7.4 (see Recipe 4)		
183					



184	C.	Co	nsumables
185		1.	Thin wall borosilicate capillaries with filament [internal radius: 0.78 mm, external radius: 1 mm],
186			used for the fabrication of the injection micropipettes (Harvard Apparatus, catalog number: 30-
187			0038)
188		2.	Thickness-selected glass coverslip (Menzel Gläser, 26 mm $\times$ 56 mm, 0.17 mm thick, custom
189			made)
190		3.	Hamilton syringe 1 mL (Carl Roth, catalog number: EY44.1)
191		4.	0.22 µm membrane filter (Milex-GV, catalog number: SLGVR04NL)
192		5.	Silicone paste (Korasilon-Paste, Carl Roth, catalog number: 0857.1)
193		6.	2 mm thick homemade Teflon spacer [length/width 40/22 mm], see Figure 1
194		7.	Eppendorf <sup>®</sup> microloader 20 μl tips (Eppendorf, catalog number: 5242956003)
195		8.	Glass vial for the lipid stock solution (DKW Life Science, catalog number: 11768929)
196		9.	Glass vial for the PVA solution (rollrandglaeser, Carl Roth, catalog number: X661.1)
197		10.	Pipette tips (capacity 1000 μl, Merk, catalog number: AXYT1000B)
198		11.	Weighing pan ROTILABO <sup>®</sup> (Roth, catalog number: 2149.2)
199		12.	Measuring vessels for Osmomat 3000 (Gonotec, catalog number: 30.9.0010)
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201	Eq	uipn	nent
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203		1.	Oven (Thermo Fisher Scientific, Heraeus Vacutherm)
204		2.	Vacuum pump (Vacuubrand, model: MZ 2C NT + 2AK)
205		3.	Micropipette puller (Sutter Instruments Model P-97)
206		4.	Micropipette manipulator system (Sutter Instruments, MPC-385)
207		5.	Microinjector (Eppendorf, FemtoJet 5247)
208		6.	Confocal microscope (Leica TCS SP5)
209		7.	Vacuum dessiccator ROTILABO <sup>®</sup> (Roth, article No. 1008.1)
210		8.	Milli-Q <sup>®</sup> system (SG water purification system, Integra UV plus, 18.2 M $\Omega$ .cm)
211		9.	Variable volume pipette (0.1 – 2 µl, 0.5 – 10 µl, 100 – 1000 µl, Eppendorf®)
212		10.	Magnetic stirrer (IKA Werke Staufen , Type: Bigsquid)
213		11.	Magnetic stirrer bars (Fisherbrand, catalog number: 11834792)
214		12.	Osmometer (Gonotec, Osmomat 3000 freezing point osmometer)
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216	Pro	oced	lure
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218	Α.	For	mation of GUVs by PVA gel-assisted swelling
219		1.	Clean a 26 mm $\times$ 56 mm glass slide by rinsing with water, ethanol, water and dry it under a $N_2$
220			flow. A separate slide is needed for each tested condition.
221		2.	Place 50 $\mu l$ of 5% w/v PVA solution (see Recipe 1) on the cleaned glass slide and spread it with
222			the micropipette tip.



223	3.	Incubate the glass at 50 °C in the oven for at least 30 min.
224	4.	Depending on the protein type to be examined, prepare a mixture of POPC:POPS:DSPE-PEG
225		2000 Biotin:DPPE-Rhodamine or POPC:POPS:chol:PI(3)P:DSPE-PEG 2000 Biotin:DPPE-
226		Rhodamine (see Recipe 2 and 3) in chloroform at 1 mg/mL final concentration.
227	5.	Clean thoroughly a Hamilton syringe with chloroform and spread evenly 10 to15 $\mu I$ of the lipid
228		mixture on the dried PVA film (taken from the oven without cooling it down) using the needle of
229		the syringe and until the slide appears dry.
230	6.	Place the glass slide for 1 h under vacuum to eliminate the excess chloroform.
231	7.	Glue the Teflon spacer (via silicone grease) to the glass with the dried lipid (see Figure 1A).
232	8.	Fill the chamber with 1800 $\mu l$ of 1× protein buffer (see Recipe 4) and place a glass coverslip on
233		top of the Teflon spacer (see Figure 1B) to avoid unwanted evaporation.
234	9.	Incubate for 10 min at room temperature to allow swelling and GUV formation.

- 235 10. After this time, tap gently a few times on the bottom of the growing chamber, remove the upper
   236 coverslip by sliding it to the side and collect the GUVs using a 1000 μl pipette tip without touching
   237 the PVA film to avoid collecting PVA debris.
- 238



- Figure 1. GUV chamber for gel-assisted swelling. The Teflon spacer is between two cover glasses (seen in panel B). The bottom glass is coated with PVA where lipids are deposited. The white oval roughly indicates the area with the deposited lipid mixture.
- B. Fabrication and loading of the micropipette
- 2451. Take a borosilicate capillary and carefully apply N2 flow through it to make sure that the capillary246is not clogged.
- 247 2. Place the capillary at the holder of the micropipette puller (Figure 2).
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### Figure 2. Capillary placed in the micropipette puller.

- 3. Pull the capillary using the one-line program to achieve a bee-needle orifice of ~250 nm (HEAT Ramp; PULL 100, VEL 10, TIME 250, PRESSURE 500') in the micropipette puller.
- 4. Filter the injection solution (2.4 µM PfBro1 and 4.8 µM of either PfVps32 or PfVps60 dissolved in 0.8× protein buffer or 0.6 µM EhVps20t and 2.4 µM EhVps32, and 0.03 mg/mL PEG-FITC to 255 256 monitor the injection; omit the addition of PEG-FITC in case one of the proteins is fluorescently 257 labelled) through the 0.22 µm filter.
  - 5. Fill the micropipette with 10 µl of the filtered protein solution using a microloader pipette tip (Figure 3).
  - 6. Tap gently the filled micropipette and leave it for 30 min in a vertical position to eliminate air bubbles.



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264 Figure 3. Loading of ESCRT proteins into the injection micropipette. The filament of the

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microloader tip is inserted as deep as possible into the injection micropipette. Pipette the loading solution into the micropipette and slowly pull out the microloader after loading the 10 µl.



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268	C.	Coat	ing of coverslip glasses with avidin and immobilization of GUVs
269		1.	Clean a 26 mm $\times$ 56 mm coverslip by rinsing with water, ethanol, water and dry it under $N_2$
270			flow.
271		2.	Functionalize the coverslips with 100 $\mu l$ of a 1:1 (v/v) mixture of BSA (1 mg/mL) and biotin-
272			BSA (1 mg/mL), dissolved in 1× protein buffer, following the protocol previously reported
273			(Maan <i>et al.</i> , 2018).
274		3.	Incubate for 20 min at room temperature.
275		4.	Rinse the glass with water and spread 100 $\mu I$ of 0.005 mg/mL avidin solution (in 1× protein
276			buffer).
277		5.	Incubate for 5 min at room temperature.
278		6.	Rinse the glass with water to remove the unbound avidin.
279		7.	Glue the Teflon spacer (via silicone paste) to the glass to form an observation chamber.
280		8.	Transfer the collected GUVs to the observation chamber with a pipette and let them sediment
281			for at least 10 min. DSPE-PEG 2000 Biotin (in the vesicle membrane) binds to the avidin on
282			the coverslip, resulting in vesicle immobilization.
283			Note: increased concentrations of avidin result in higher binding of the GUV to the glass
284			surface which later hinders budding as the vesicle excess area is consumed for adhesion
285			(see Figure 4).
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Figure 4. Side view of immobilized vesicles (here, POPC:POPS:chol:Pl(3)P:DSPE-PEG
 2000 Biotin:DPPE-Rhodamine in molar ratio 60.9:10:25:3:1:0.1). The coverslips were
 coated with a 1:1 mixture of biotin-BSA (1 mg/mL) and BSA (1 mg/mL), and then avidin was
 deposited. The vesicles are visualized via vertical confocal cross sections. The color is inverted.
 The scale bars correspond to 10 µm.



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- 294 D. Injection and observation of GUVs
  - 1. Place the observation chamber with the immobilized GUVs on the microscope stage.
  - 2. Connect the filled micropipette to the holder of the microinjector (Figure 5).



Micromanipulator

Micropipette holder of the micromanipulator

Micropipette holder of the microinjector

Figure 5. Attachment of the micropipette holder of the microinjector to the micromanipulator.

- 3. Attach the holder with the pipette to the mechanical arm of the micromanipulator.
- 4. Set the angle of injection as large as the microscope setup allows (see Figure 6).
- Note: The most ideal angle of injection would be 90° to avoid lateral displacement of the vesicles during puncture attempts, but such cannot be achieved since the condenser of the microscope occupies the space above the sample.



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Figure 6. Micropipette setup of the injection procedure. Panel A: top-view of the setup; for clarity the observation chamber was removed. Panel B: a close-up side-view of the injection angle. Note that the micropipette holder is as close to the condenser of the microscope as possible to ensure high angle of injection.



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313	5.	With the micromanipulator, introduce the micropipette into the solution of the observation
314		chamber and focus on the tip of the pipette.
315	6.	Set the microscope to the desired observation settings. For this measurement they were the
316		following:
317		- The DPPE-Rhodamine dye (integrated in the membrane of the GUVs) was excited with
318		a diode-pumped solid-state 561 nm laser and the signal was collected in the range 570-
319		650 nm.
320		- The PEG-FITC dye was excited with a 488 nM line of an argon laser and the signal was
321		collected in the range 495-530 nm.
322		- To avoid crosstalk between the different fluorescence signals, sequential scanning was
323		performed.
324	7.	Place the micropipette in a site where no GUVs are observed and purge it (by pressing the
325		"Clean" bottom of the microinjector) to confirm that the micropipette is not clogged; a signal
326		from the fluorescent dye leaving the pipette tip should be detected.
327	8.	Lower the micropipette close to but still above the focal plane of the GUVs.
328	9.	Approach a selected GUV with the tip of the micropipette from above; the selected vesicle
329		should be clean and without defects (to ensure this, examine the selected GUV with a XYZ
330		scan).
331	10	. Lift the micropipette a few micrometers above the vesicle (the micropipette tip goes out of
332		focus).
333	11	. Puncture the GUV by moving the micropipette towards the vesicle in both Z and X directions.
334	12	. Perform a XYZ scan to make sure that the micropipette penetrated into the GUV.
335	13	. Start recording a time sequence.
336	14	. Inject the vesicle (see Figure 7 and Figure 8) using the following parameters of the
337		microinjector: pressure of injection = 150 hPa; time of injection = 5 s; compensation pressure
338		= 1 hPa. These parameters ensure injecting volumes in sub-picoliter range (a few hundreds
339		of femtoliters).
340	15	. Pull the micropipette out from the interior of the GUV (lift the micropipette in Z direction).
341	16	. Perform a XYZ scan to detect in which focal plane the outward buds formed (Figure 7, Figure
342		8).
343		Note: tense vesicles are easier to puncture than fluctuating ones.
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Figure 8. Horizontal (xy) confocal cross sections of injection of GUVs with PEG-FITC solution (top row) or *Entamoeba histolytica* purified ESCRT-III proteins (bottom row). The membrane is shown in magenta and PEG-FITC in cyan. The tip of the injection pipette can be seen on the first two frames in each sequence. The arrows on the last frame point to the outward buds; see also supplementary <u>Movie S2</u>. The inset in the last snapshot shows zoomed-in view of outward buds from the same vesicle but at another xy plane.



364	<u>Recipe</u>	<u>es</u>						
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366	1.	PVA s	olution (5%	5 <b>w/v)</b>				
367		a. W	/eigh 0.5 g F	PVA and p	lace it in a glass vi	al.		
368		b. A	dd 10 mL of	<sup>1</sup> × protei	n buffer (to mainta	in osmolarity) a	and place a clean magnetic	stirrer
369		in	the vial.					
370		c. K	eep the sol	ution in a	90 °C water bath	under constan	t stirring (300-400 rpm) uni	iil PVA
371		di	issolves con	npletely. T	he solution becom	es clear.		
372	2.	Lipid	mix (1 mg/	mL) for th	ne injection of ES	CRTs purified	l from <i>P. falciparum</i> cont	aining
373		POPC	:POPS:DSP	PE-PEG 2	000 Biotin:DPPE-	Rhodamine (F	inal volume 200 μl)	
		Lipi	ds	(stock	Concentration	Volume (µl)	Final molar fraction %	
		solu	utions) & R	eagents	(mg/mL)			
		POF	PC		10	15.78	78.9	
		POF	PS		10	4.00	20.0	

0.25

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375 376 377  Lipid mix (1 mg/mL) for the injection of ESCRTs purified from *Entamoeba histolytica* containing POPC:POPS:chol:PI(3)P:DSPE-PEG 2000 Biotin:DPPE-Rhodamine (Final volume 200 μl)

2.00

1.00

177.22

1.0

0.1

Lipids (stock solutions) &	Concentration	Volume (µl)	Final molar
Reagents	(mg/mL)		fraction %
POPC	10	12.18	60.9
POPS	10	2.00	10
chol	10	5.00	25
PI(3)P (dissolved in	0.25	24	3
chloroform/methanol in 60 to 40 ratio)			
DSPE-PEG 2000 Biotin	1	2.00	1.0
DPPE-Rhodamine	0.25	1.00	0.1
Chloroform		153.82	

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## **4. Protein buffer 1×, pH 7.4**

DSPE-PEG 2000 Biotin

DPPE-Rhodamine

Chloroform

Compound	Concentration (mM)	Osmolarity (mOsm)
Trizma® hydrochloride	25	325
NaCl	150	
Adjust pH with 5 M HCI		



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#### 395 Competing interests

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397 The authors declare no competing interests.

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